

OIPE

RAW SEQUENCE LISTING

DATE: 12/20/2001

PATENT APPLICATION: US/09/771,383

TIME: 14:15:05

Input Set : A:\2368-12.ST25.txt

Output Set: N:\CRF3\12202001\I771383.raw

ENTERED

3 <110> APPLICANT: Parris, Kevin
4 Somers, William
5 Tam, Amy
6 Lin, Laura
7 Stahl, Mark
9 <120> TITLE OF INVENTION: CRYSTAL STRUCTURE OF ACYL CARRIER PROTEIN SYNTHASE
10 AND ACYL CARRIER PROTEIN SYNTHASE COMPLEX
12 <130> FILE REFERENCE: 2368/12
14 <140> CURRENT APPLICATION NUMBER: US 09/771,383
16 <141> CURRENT FILING DATE: 2001-01-25
18 <150> PRIOR APPLICATION NUMBER: US 60/178,639
20 <151> PRIOR FILING DATE: 2000-01-28
22 <160> NUMBER OF SEQ ID NOS: 13
24 <170> SOFTWARE: PatentIn version 3.0
26 <210> SEQ ID NO: 1
28 <211> LENGTH: 121
30 <212> TYPE: PRT
32 <213> ORGANISM: B. subtilis
34 <400> SEQUENCE: 1
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39 Ser Met Ala Gly Arg Gln Lys Arg Phe Ala Glu Arg Ile Leu Thr Arg
40 20 25 30
42 Ser Glu Leu Asp Gln Tyr Tyr Glu Leu Ser Glu Lys Arg Lys Asn Glu
43 35 40 45
45 Phe Leu Ala Gly Arg Phe Ala Ala Lys Glu Ala Phe Ser Lys Ala Phe
46 50 55 60
48 Gly Thr Gly Ile Gly Arg Gln Leu Ser Phe Gln Asp Ile Glu Ile Arg
49 65 70 75 80
51 Lys Asp Gln Asn Gly Lys Pro Tyr Ile Ile Cys Thr Lys Leu Ser Gln
52 85 90 95
54 Ala Ala Val His Val Ser Ile Thr His Thr Lys Glu Tyr Ala Ala Ala
55 100 105 110
57 Gln Val Val Ile Glu Arg Leu Ser Ser
58 115 120
61 <210> SEQ ID NO: 2
63 <211> LENGTH: 122
65 <212> TYPE: PRT
67 <213> ORGANISM: Aquifex
69 <400> SEQUENCE: 2
71 Met Ile Gly Val Asp Ile Val Lys Asn Glu Arg Ile Lys Asp Ala Leu
72 1 5 10 15
74 Glu Arg Phe Gly Asp Lys Phe Leu Asp Arg Ile Tyr Thr Lys Arg Glu
75 20 25 30
77 Leu Glu Tyr Cys Tyr Ala His Cys Asp Phe Leu Pro Cys Leu Ala Ala
78 35 40 45
80 Arg Trp Ala Gly Lys Glu Ala Val Leu Lys Ala Phe Tyr Thr Glu Phe

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81      50      55      60
83 Lys Ile Phe Leu Arg Phe Lys Glu Ile Glu Ile Leu Gly Asn Arg Gly
84 65      70      75      80
86 Arg Pro Pro Thr Val Ile Asn Arg Glu Gly Val Glu Glu Ile Leu
87      85      90      95
89 Lys Asn Tyr Glu Val Ile Val Ser Leu Ser His Glu Arg Asp Tyr Ser
90      100      105      110
92 Val Ala Val Ala Tyr Ile Lys Lys Lys Ser
93      115      120
96 <210> SEQ ID NO: 3
98 <211> LENGTH: 122
100 <212> TYPE: PRT
102 <213> ORGANISM: Chlamydophila
104 <400> SEQUENCE: 3
106 Met Glu Ile Ile His Ile Gly Thr Asp Ile Ile Glu Ile Ser Arg Ile
107 1      5      10      15
109 Arg Glu Ala Ile Ala Thr His Gly Asn Arg Leu Leu Asn Arg Ile Phe
110      20      25      30
112 Thr Glu Ala Glu Gln Lys Tyr Cys Leu Glu Lys Thr Asp Pro Ile Pro
113      35      40      45
115 Ser Phe Ala Gly Arg Phe Ala Gly Lys Glu Ala Val Ala Lys Ala Leu
116      50      55      60
118 Gly Thr Gly Ile Gly Ser Val Val Ala Trp Lys Asp Ile Glu Val Phe
119 65      70      75      80
121 Lys Val Ser His Gly Pro Glu Val Leu Leu Pro Ser His Val Tyr Ala
122      85      90      95
124 Lys Ile Gly Ile Ser Lys Val Ile Leu Ser Ile Ser His Cys Lys Glu
125      100      105      110
127 Tyr Ala Thr Ala Thr Ala Ile Ala Leu Ala
128      115      120
131 <210> SEQ ID NO: 4
133 <211> LENGTH: 119
135 <212> TYPE: PRT
137 <213> ORGANISM: Helicobacter
139 <400> SEQUENCE: 4
141 Met Ile Gly Ile Asp Ile Val Ser Ile Ala Arg Ile Glu Lys Cys Val
142 1      5      10      15
144 Lys Arg Phe Lys Met Lys Phe Leu Glu Arg Phe Leu Ser Pro Ser Glu
145      20      25      30
147 Ile Val Leu Cys Lys Asp Lys Ser Ser Ile Ala Gly Phe Phe Ala
148      35      40      45
150 Leu Lys Glu Ala Cys Ser Lys Ala Leu Gln Val Gly Ile Gly Lys Glu
151      50      55      60
153 Leu Ser Phe Leu Asp Ile Lys Ile Ser Lys Ser Pro Lys Asn Ala Pro
154 65      70      75      80
156 Leu Ile Thr Leu Ser Lys Glu Lys Met Asp Tyr Phe Asn Ile Gln Ser
157      85      90      95
159 Leu Ser Ala Ser Ile Ser His Asp Ala Gly Phe Ala Ile Ala Val Val
160      100      105      110

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162 Val Val Ser Ser Ser Asn Glu
163      115
166 <210> SEQ ID NO: 5
168 <211> LENGTH: 119
170 <212> TYPE: PRT
172 <213> ORGANISM: Staphylococcus
174 <400> SEQUENCE: 5
176 Met Ile His Gly Ile Gly Val Asp Leu Ile Glu Ile Asp Arg Ile Gln
177 1      5      10      15
179 Ala Leu Tyr Ser Lys Gln Pro Lys Leu Val Glu Arg Ile Leu Thr Lys
180      20      25      30
182 Asn Glu Gln His Lys Phe Asn Asn Phe Thr His Glu Gln Arg Lys Ile
183      35      40      45
185 Glu Phe Leu Ala Gly Arg Phe Ala Thr Lys Glu Ala Phe Ser Lys Ala
186      50      55      60
188 Leu Gly Thr Gly Leu Gly Lys His Val Ala Phe Asn Asp Ile Asp Cys
189 65      70      75      80
191 Tyr Asn Asp Glu Leu Gly Lys Pro Lys Ile Asp Tyr Glu Gly Phe Ile
192      85      90      95
194 Val His Val Ser Ile Ser His Thr Glu His Tyr Ala Met Ser Gln Val
195      100     105     110
197 Val Leu Glu Lys Ser Ala Phe
198      115
201 <210> SEQ ID NO: 6
203 <211> LENGTH: 169
205 <212> TYPE: PRT
207 <213> ORGANISM: Thermotoga
209 <400> SEQUENCE: 6
211 Met Ile Val Gly Val Gly Ile Asp Val Leu Glu Val Glu Arg Val Pro
212 1      5      10      15
214 Glu Lys Phe Ala Glu Arg Ile Leu Gly Glu Ser Glu Lys Arg Leu Phe
215      20      25      30
217 Leu Thr Arg Lys Arg Arg Arg Glu Phe Ile Ala Gly Arg Phe Ala Leu
218      35      40      45
220 Lys Glu Ala Phe Phe Lys Ala Leu Gly Thr Gly Leu Asn Gly His Ser
221      50      55      60
223 Phe Thr Asp Val Glu Phe Leu Glu Ser Asn Gly Lys Pro Val Leu Cys
224 65      70      75      80
226 Val His Lys Asp Phe Gly Phe Phe Asn Tyr Ala His Val Ser Leu Ser
227      85      90      95
229 His Asp Arg Phe Ala Val Ala Leu Val Val Leu Glu Lys Arg Lys Gly
230      100     105     110
232 Asp Ile Ile Val Glu Gly Asp Glu Ser Phe Leu Arg Lys Arg Phe Glu
233      115     120     125
235 Val Leu Glu Arg Ser Val Glu Gly Trp Glu Ile Glu Thr Ser Leu Pro
236      130     135     140
238 Pro Phe Thr Leu Lys Lys Leu Leu Glu Ser Ser Gly Cys Arg Leu Val
239 145     150     155     160
241 Arg Tyr Gly Asn Ile Leu Ile Gly Glu

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242                               165
245 <210> SEQ ID NO: 7
247 <211> LENGTH: 126
249 <212> TYPE: PRT
251 <213> ORGANISM: Escherichia
253 <400> SEQUENCE: 7
255 Met Ala Ile Leu Gly Leu Gly Thr Asp Ile Val Glu Ile Ala Arg Ile
256 1                               5                10                15
258 Glu Ala Val Ile Ala Arg Ser Gly Asp Arg Leu Ala Arg Arg Val Leu
259                               20                25                30
261 Ser Asp Asn Glu Trp Ala Ile Trp Lys Thr His His Gln Pro Val Arg
262                               35                40                45
264 Phe Leu Ala Lys Arg Phe Ala Val Lys Glu Ala Ala Ala Lys Ala Phe
265                               50                55                60
267 Gly Thr Gly Ile Arg Asn Gly Leu Ala Phe Asn Gln Phe Glu Val Phe
268 65                               70                75                80
270 Asn Asp Glu Leu Gly Lys Pro Arg Leu Arg Leu Trp Gly Glu Ala Leu
271                               85                90                95
273 Lys Leu Ala Glu Lys Leu Gly Val Ala Asn Met His Val Thr Leu Ala
274                               100               105               110
276 Asp Glu Arg His Tyr Ala Cys Ala Thr Val Ile Ile Glu Ser
277                               115               120               125
280 <210> SEQ ID NO: 8
282 <211> LENGTH: 126
284 <212> TYPE: PRT
286 <213> ORGANISM: Rickettsia
288 <400> SEQUENCE: 8
290 Met Leu Ile Gly Val Gly Thr Asp Ile Val Gln Ile Pro Arg Ile Glu
291 1                               5                10                15
293 Lys Ile Leu Asn Ile Tyr Gln Glu Leu Phe Ala Lys Lys Ile Leu Ala
294                               20                25                30
296 Leu Lys Glu Leu Lys Gln Phe Thr Leu Leu Asn Lys Thr Asn His Ala
297                               35                40                45
299 Thr Phe Leu Ala Lys Arg Phe Ser Ala Lys Glu Ala Val Ser Lys Ala
300                               50                55                60
302 Phe Gly Val Gly Ile Gly Arg Gly Ile Asn Phe Lys Asp Ile Thr Ile
303 65                               70                75                80
305 Leu Asn Asp Asn Leu Gly Lys Pro Thr Val Glu Ile Ser Ser His Tyr
306                               85                90                95
308 Thr Asn Lys Leu Ala Pro Phe Asn Ile His Leu Ser Leu Ser Asp Asp
309                               100               105               110
311 Tyr Pro Ile Cys Ile Ala Phe Ala Ile Ile Glu Ser Asn Cys
312                               115               120               125
315 <210> SEQ ID NO: 9
317 <211> LENGTH: 123
319 <212> TYPE: PRT
321 <213> ORGANISM: Streptomyces
323 <400> SEQUENCE: 9
326 Met Ser Ile Ile Gly Val Gly Ile Asp Val Ala Glu Val Glu Arg Phe

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327 1          5          10          15
329 Gly Ala Ala Leu Glu Arg Thr Pro Ala Leu Ala Gly Arg Leu Phe Leu
330          20          25          30
332 Glu Ser Glu Leu Leu Leu Pro Gly Glu Arg Arg Gly Val Ala Ser
333          35          40          45
335 Leu Ala Ala Arg Phe Ala Ala Lys Glu Ala Leu Ala Lys Ala Leu Gly
336          50          55          60
338 Ala Pro Ala Gly Leu Leu Trp Thr Asp Ala Glu Val Trp Val Glu Ala
339 65          70          75          80
341 Gly Gly Arg Pro Arg Leu Arg Val Thr Gly Thr Val Ala Ala Arg Ala
342          85          90          95
344 Ala Glu Leu Gly Val Ala Ser Trp His Val Ser Leu Ser His Asp Ala
345          100          105          110
347 Gly Ile Ala Ser Ala Val Val Ile Ala Glu Gly
348          115          120
351 <210> SEQ ID NO: 10
353 <211> LENGTH: 125
355 <212> TYPE: PRT
357 <213> ORGANISM: Treponema
359 <400> SEQUENCE: 10
361 Met Ile Ile Gly Val Gly Ile Asp Ile Val Glu Ile Glu Arg Phe Val
362 1          5          10          15
364 Ser Trp Thr His Asn Val Arg Leu Leu Arg Arg Phe Phe His Gln Glu
365          20          25          30
367 Glu Ile Val Asp Phe Phe Lys Asn His Met Arg Ala Gln Phe Leu Ala
368          35          40          45
370 Thr Arg Phe Ala Ala Lys Glu Ala Phe Gly Lys Ala Leu Gly Thr Gly
371          50          55          60
373 Leu Arg Asn Met Glu Leu Arg Asn Ile Arg Val Cys Gln Asn Gly Trp
374 65          70          75          80
376 Gly Lys Pro Arg Leu Glu Val Tyr Gly Ala Ala Gln Ala Met Leu Ala
377          85          90          95
379 Ala Thr Gly Gly Thr His Ile Gln Val Ser Leu Thr His Glu Arg Glu
380          100          105          110
382 Val Ala Ser Ala Ile Val Ile Ile Glu Gly Glu Pro Leu
383          115          120          125
386 <210> SEQ ID NO: 11
388 <211> LENGTH: 121
390 <212> TYPE: PRT
392 <213> ORGANISM: Bacillus
394 <400> SEQUENCE: 11
396 Met Ile Tyr Gly Ile Gly Leu Asp Ile Thr Glu Leu Lys Arg Ile Ala
397 1          5          10          15
399 Ser Met Ala Gly Arg Gln Lys Arg Phe Ala Glu Arg Ile Leu Thr Arg
400          20          25          30
402 Ser Glu Leu Asp Gln Tyr Tyr Glu Leu Ser Glu Lys Arg Lys Asn Glu
403          35          40          45
405 Phe Leu Ala Gly Arg Phe Ala Ala Lys Glu Ala Phe Ser Lys Ala Phe
406          50          55          60

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VERIFICATION SUMMARY

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Input Set : A:\2368-12.ST25.txt

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